Search Results

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E-VALUE COLOR KEY

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WU-BLAST 2.0MP-WashU [05-May-2004] [irix6.5-r10k-132LPF64 2004-05-<= E-75 stronger 06T14:17:11] huFlt3-L vs database nr_aa (702746936 letters in 2091158 sequences) E-30 - E-75 SIMILARITY E-5 - E-30 1.0 - 2-5 Forwards selected sequences to another application. > 1.0 weaker Query= huflt3-L 28-160 Alignment vs. Query (Click to View Alignment) (length=133) 100 50 Sequences producing Righ-scoring Segment Score E-Value Pairs: pir | 139076 F1t3 D ligand alternatively spliced isoform -_____100%<u>____</u> human >gb|A.. ref NP_001450.2 fmsrelated tyrosine kinase 3 ligand [Homo 701 2.2E-68 ----sapiens.. pdb | 1ETED Chain D. Crystal Structure Of The Flt3 Ligand -99% 698 4.5R-68 ----->pdb | 1ETE | ... rf|2009389B prf|20093K9B ligand -99%-695 9.4E-68 ----pir | S43293 FLT3/FLK2 --99%-1 fligand (clone \$109) human ☐ **6** gb|AAA17999.1 FLT3/FLK2 ligand 695 9.4E-68 ______99%-gb|AA072538.1 FLT3 1 6 ligand [Papio 683 1.8E-66 -----97%---cynocephalus x Papio anubis] ☐ **6** gb|AAF87089.1 Flt3 ligand [Felis catus] 581 1.1E-55 _-----84%ref|NP_001003350.1 Fit3 ligand [Canis familiaris] 581 1.1E-55 _-------84<u>%----</u> >gb|AAF87068.1| .. dbj|BAB79634.1 flt3 ______81%------569 2.1E-54 ----ref NP_851373.1 fmsrelated tyrosine kinase 3 ligand [Bos 569 2.1E-54 ----taurus] .. gb|AAA90952.1 Flt3 508 6.2E-48 _______73%--1 1 ligand alternatively spliced isoform pir|158343 flt3 mouse >pir | | 543291

☐ **1** gb[AAD38036.1 tWs opsin [Columba livia]

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Search Results	
FLT3/FLK2 1	
gb AAA93305.1 Flt3 ligand, Tl69 form	508 6.22-48>
sp P49772 SL cytokine precursor (Fms- related tyrosine kinase 3 lig.	508 6.2E-48>
pir A49265 flt3/flk-2 ligand precursor - mouse >gb AAA39436.1 flt	508 6.2E-48>
ref NP_038548.2 FMS- like tyrosine kinase 3 ligand [Mus musculus] >	504 1.68-47
ref XP_341855.1 similar to RIKEN cDNA 2410004H02 [Rattus norvegicu	494 7.1E-46
gb AAF99323.1 flt3 fligand isoform-2 (Bostaurus)	369 3.4E-45
gb AAH19801.1 Flt31 protein (Mus musculus)	481 4.5E-45>
gb AAH06331.1 FLT3LG protein [Homo sapiens] >gb AAH11914.1 FLT3LG	276 2.4E-23100%
emb CAA48516.1 myocyte-specific enhancer factor 2 (aMEF2) [Homo sa	59 3.7 <u>41%></u>
ref[NP_269539.1 hypothetical protein _ phage associated [streptoco	.59 3.7 <u>45%></u>
gene product	58 4.0 <u>32%></u>